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(54) Title: PHARMACEUTICAL LIPID AGGREGATE WITH HELICOBACTER PYLORI ANTIGEN AND NEGATIVELY CHARGED LIPID

#### (57) Abstract

The present invention relates to a new pharmaceutical formulation comprising a lipid aggregate of a negatively charged lipid or lipid mixture and at least one antigenic, native or recombinant polypeptide which constitute a *Helicobacter pylori* antigen, and optionally a pharmaceutically acceptable carrier. More specifically the invention relates to a lipid aggregate comprising a surface–exposed antigen with an approximate molecular weight of 29 kDa. Furthermore the invention provides a formulation of a nucleic acid molecule coding for the said polypeptide. The new formulations, which are useful as vaccine formulations, elicit a protective immune response against *H. pylori* infections, and said vaccine formulations are suitable for both therapeutic and prophylatic use.

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PHARMACEUTICAL LIPID AGGREGATE WITH HELICOBACTER PYLORI ANTIGEN AND NEGATIVELY CHARGED LIPID

#### Field of the invention

The present invention provides a new pharmaceutical formulation comprising polypeptides, which constitute *Helicobacter pylori* antigens, or nucleic acid molecules coding for the said polypeptides. More specifically the present invention relates to a pharmaceutical formulation in the form of a lipid aggregate comprising these polypeptides or antigens. Furthermore, the invention provides a process for the preparation of such a pharmaceutical formulation, and the use of such a formulation in the treatment of *Helicobacter pylori* infections, as well as a method of treating a patient to elicit a protective immune response against *Helicobacter pylori* infections, said formulation being suitable for both therapeutic and prophylactic use.

#### 15 Background of the invention

There is a need for an effective drug delivery system, such as a vaccine composition, for polypeptides or antigens involved in the protection and treatment of *H. pylori* infections. The pharmaceutical formulation shall provide an improved immune response to the polypeptides or antigens delivered by the formulation.

The gram-negative bacterium *Helicobacter pylori* is an important human pathogen, involved in several gastroduodenal diseases. Colonization of gastric epithelium by the bacterium leads to active inflammation and progressive chronic gastritis, with a greatly enhanced risk of progression to peptic ulcer disease.

In order to colonize the gastric mucosa, *H. pylori* uses a number of virulence factors. Such virulence factors comprise several adhesins, with which the bacterium associates with the mucus and/or binds to epithelial cells; ureases which helps to neutralize the acid environment; and proteolytic enzymes which makes the mucus more fluid.

Despite a strong apparent host immune response to *H. pylori*, with production of both local (mucosal) as well as systemic antibodies, the pathogen persists in the gastric mucosa, normally for the life of the host. The reason for this is probably that the spontaneously induced immune-response is inadequate or directed towards the wrong epitopes of the antigens.

In order to understand the pathogenesis and immunology of *H. pylori* infections, it is of great importance to define the antigenic structure of this bacterium. In particular, there is a need for characterization of surface-exposed (like adhesins) and secreted proteins which, in many bacterial pathogens, have been shown to constitute the main virulence factors, and which can be useful for the diagnosis of *H. Pylori* and in the manufacture of vaccine compositions.

- Cloning of the *hpaA* gene, which codes for a 20 kDa receptor-binding subunit of the *N*-acetylneuraminyllactose-binding fibrillar hemagglutinin (NLBH) of *H. pylori*, has been disclosed by Evans et al. (1993) J. Bacteriol. 175, 674-683.
- Monoclonal antibodies (MAbs) against membrane preparations of *H. pylori* have been disclosed by Bölin et al. (1995) J. Clin. Microbiol. 33, 381-384. One of these MAbs, designated HP30-1:1:6, reacted with a 30 kDa protein which was shown to be exposed on the surface of intact bacteria and to have properties like that of an adhesin.
- Whenever stressed or threatened, the *H. pylori* cell transforms from a bacillary to a coccoid form. In the coccoid form, the *H. pylori* cell is much less sensitive to antibiotics and other anti-bacterial agents. Circumstantial evidence indicate the *H. pylori* might be transmitted between individuals in this form, possibly via water or direct contact. An efficient vaccine composition should therefore elicit an immune response towards both the coccoid and the bacillary form of *H. pylori*. Since systemic immunity probably only plays a limited role in protection against mucosal infections,

it is also important that the vaccine composition will enhance protective immune mechanisms locally in the stomach.

### Prior Art

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Several different types of vaccine formulations have been described in the literature (see e g "Vaccine Design. The subunit and adjuvant approach" (Eds: M F Powell and M J Newman), Pharmaceutical Biotechnology, vol 6, Plenum Press, NY 1995). Thus, vaccines have been formulated in known delivery systems such as liposomes,

ISCOMs, cochleates, etc, or have been attached to or included into polymer microspheres of degradable or non-degradable nature. Antigens have been associated with live attenuated bacteria, viruses or phages or with killed vectors of the same kind. Other acceptable carriers or diluents, known to those skilled in the art, are e g phosphate buffered saline, enterically coated powder formulations, surface active substances and polymers, etc.

Specifically, aggregates formed between amphiphilic molecules, e g phospholipids, and ions of opposite charge, have been studied extensively (e g Nir et al, Progress in Surface Science, 13 (1983), 1 - 124) and also sometimes been reported as suitable formulations or formulation components for different types of therapeutic agents, including antigens. In some of these reports the focus have been on the use of negatively charged phospholipids, e g phoshatidylserine, phosphatidylglycerol, etc, and positively charged ions like Ca<sup>2+</sup>, Mg<sup>2+</sup>, etc. Large, cylindrical structures with lipid multilayers in a spiral configuration was reported to exist in mixtures of phosphatidylserine and Ca<sup>2+</sup> in water by Papahadjopoulos et al (*Biochim Biophys Acta* 394, 483-491 (1975)) and named cochleates by the authors. The morphology of these structures as well as prerequisites and techniques for their formation, either in pure phospholipid systems, or in combination with other substances, e g other phospholipids, cholesterol, polypeptides like antigens, DNA etc, have been reported in a number of studies (e g Galla and Sackmann, *Biochim Biophys Acta* 401, 509-529 (1975); Papahadjopolous et al. *Biochim Biophys Acta* 465, 579-598 (1977); Wilschut and

Papahadjopoulus Nature 281, 690-692 (1979); Tilcock et al. Biochemistry 23, 2696-2703 (1984); Silvius and Gagné Biochemistry 23, 3241-3247 (1984); Graham et al. Biochemistry 24, 7123-7131 (1984); Gould-Fogerite and Mannino Analytical Biochemistry 148, 15-25 (1985) Kouaouci et al. Biochemistry 24, 7132-7140 (1985); Gould-Fogerite et al. in Advances in Membrane Biochemistry and Bioenergetics (Eds: Kim et al.), Plenum Press NY 1988, pp 569-586; Mannino and Gould-Fogerite *Biotechniques* <u>6</u>, 682-690 (1988); Gould-Fogerite et al. Gene <u>84</u>, 429-438 (1989); Roux and Bloom *Biophys J* <u>60</u>, 38-44 (1991); Flach and Mendelsohn Biophys J 64, 1113-1121 (1993); Gould-Fogerite and Mannino in Liposome Technology (Ed Gregoriadis), CRC Press, Boca Raton 1993, vol I pp 67-80, and vol III pp 261-276; Nieva et al. Biochemistry 33, 3201-3209 (1994); Hinderliter et al. Biophys J 67, 1906-1911 (1994)). A few patents and patent applications have also described drug delivery systems based on either cochleate structures directly, or on systems derived from cochleate structures (Papahadjopoulos US78/4078052; Mannino and Gould-Fogerite US87/4663161; Mannino and Gould-Fogerite US89/4871488; Gould-Fogerite and Mannino WO95/09648).

#### Purpose of the invention

The purpose of this invention is to provide a new pharmaceutical formulation of an antigenic *Helicobacter pylori* polypeptide or a pharmaceutical formulation of a nucleic acid molecule coding for such a polypeptide. More specifically, the present invention relates to a new pharmaceutical formulation based on a lipid aggregate of a negatively charged lipid or mixture of lipids, and at least one polypeptide which is a Helicobacter pylori antigen, or nucleic acid molecules coding for such polypeptides, and optionally a pharmaceutically acceptable carrier. One suitable antigen for the lipid aggregate is a 29 kDa polypeptide, obtained by the recombinant cloning of a *H. pylori* gene which encodes a surface exposed protein. The 29 kDa polypeptide is described below as well as in the Applicant's co-pending patent application

WO96/38475, published 5 December 1996. The nucleic acid sequence of the 29 kDa

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polypeptide is similar to the sequence of the *hpaA* gene as published by Evans et al. (1993) in Journal of Bacteriology, vol 175, 674-683. Thus, the pharmaceutical formulation according to the present invention is also suitable for the *hpaA* gene.

In general terms, the new pharmaceutical formulation is also suitable for other *H. pylori* antigens, especially membrane proteins. In the following such suitable antigens for the formulation are the 29 kDa polypeptide described below, or the *hpaA* gene described by Evans et al. The different names 29 kDa and *hpaA* gene or HpaA protein are used interchangable in the present specification as suitable antigens for the present formulation, even if the sequence of the *hpaA* gene from different *H. pylori* strains shows some differences (P.W. Toole et al, Bacteriology Vol. 177, No. 21, Nov. 1995).

The 29 kDa polypeptide is described in the Applicant's co-pending WO96/38475, published 5 December 1996, as an antigenic protein which is expressed in all strains of *H. pylori*, also in coccoid forms of the bacterium, and which is able to induce a mucosal as well as a systemic immune-response in a host measured as antibody production. The 29 kDa polypeptide is expressed by all *H. pylori* strains tested, and antibodies created towards this protein do not cross-react with common endogenous human bacteria of other species or with selected human tissues including the gastric mucosa. Thus being an essential, well conserved adhesin with immunogenic properties, the 29 kDa polypeptide is useful both for the detection of *H. pylori* infections as well as for the manufacture of vaccine compositions. When given in an appropriate pharmaceutical formulation, as the new formulation according to the present invention, the 29 kDa polypeptide elicits a protective or therapeutic immune response against such *H. pylori* infections.

# Description of the 29 kDa polypeptide

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The 29 kDa polypeptide is described in WO96/38475 hereby incorporated in whole by reference. It has an amino sequence substantially similar to a *Helicobacter pylori* 

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surface-exposed antigen and has an approximate molecular weight of 29 kDa. The said surface-exposed antigen has i.a. the following important properties:

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- It is an adhesin, which is important for the colonization of the gastric mucosa:
- It is expressed on the surface of both dividing (bacillary) forms as well as resting (coccoid) forms of *H. pylori*;
  - It is a strong antigen giving rise to both systemic and local (mucosal) production
    of antibodies;
  - It is conserved in all tested strains of H. pylori;

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- Antibodies to the 29 kDa polypeptide do not cross-react with a number of different non-helicobacter bacteria, or with selected human tissues, including the gastric mucosa;
  - The 29 kDa polypeptide is lipidated and thus post-translationally modified. This
    feature of the polypeptide may be of importance for its immunogenicity and
    for its proper exposure on the surface of *H. pylori*. It is known in the art that
    lipid modification can be essential for the immunological properties of
    bacterial lipoproteins (see Weis, J.J. et al. (1994) Infection and Immunity, vol.
    62, 4632-4636).
  - It is a putative virulence-factor, whereby the term "virulence factor" is to be understood a molecule specifically involved in adherence of *H. pylori* to the epithelial surface of the gastric mucosa and / or in the establishment and maintenance of *H. pylori* infection.

In a preferred form, the said polypeptide has an amino acid sequence according to positions 1-260, or 28-260, in SEQ ID NO: 2 or 4 of the Sequence Listing. Further, it is believed that positions 1-260 in SEQ ID NO: 2 and 4 represent the uncleaved protein, while positions 1-27 represent a signal sequence and positions 28-260 represent the mature polypeptide. The only difference between SEQ ID NO: 2 and SEQ ID NO: 4 is that SEQ ID NO: 2 has a Ser residue in position 222, while SEQ ID NO: 4 has an Arg residue in the same position.

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However, the polypeptides which can be formulated according to the invention is not to be limited strictly to the 29 kDa polypeptide with an amino acid sequence identical with the above mentioned positions in SEQ ID NO: 2 or 4 in the Sequence Listing.

Rather the invention encompasses polypeptides carrying modifications like substitutions, small deletions, insertions or inversions, which polypeptides nevertheless have substantially the properties of the 29 kDa polypeptide. Such properties include the ability to elicit a mucosal as well as systemic immune-response against *H. pylori* in a mammal host; the ability to work as an adhesin; and the presence of the polypeptide in both bacillary and coccoid forms of *H. pylori*.

Consequently, one purpose of the invention is to present a formulation based on a lipid aggegate of polypeptides, the amino acid sequence of which is at least 90% homologous, preferably at least 95% homologous, with the amino acid sequence shown as positions 1-260, or positions 28-260, in SEQ ID NO: 2 or 4, in the Sequence Listing, which polypeptides nevertheless have substantially the biological activities of the 29 kDa polypeptide.

Further, the purpose of the invention is to provide a formulation based on a lipid aggregate of peptides, with a length of at least 5 amino acids, which comprise an immunogenic epitope of the 29 kDa polypeptide and retains the ability to elicit an immune response against *H. pylori* bacteria in a mammal host. Such epitope(s) can be presented alone or in the form of fusion proteins, where the epitope is fused to an inert or immunologically active carrier polypeptide. The identification of these epitopes will be based on the presence of host-generated antibodies towards different segments of the 29 kDa polypeptide.

One way of obtaining structural information on the epitopes of the 29 kDa polypeptide is the production and characterisation of monoclonal antibodies binding to the polypeptide, followed by mapping of epitopes by e.g. Pepscan analysis.

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Monoclonal antibodies can be produced by standard methods, such as those described by De St. Groth (1980) in J. Immunol. Methods, vol. 35, 1-21.

In another aspect, the invention provides a formulation based on a lipid aggregate of an isolated and purified nucleic acid molecule which has a nucleotide sequence coding for a polypeptide as defined above. In a preferred form of the invention, the said nucleic acid molecule is a DNA molecule which has a nucleotide sequence identical with SEQ ID NO: 1 or 3 of the Sequence Listing. However, the DNA molecule is not to be limited strictly to the sequence shown as SEQ ID NO: 1 or 3. Rather the invention encompasses formulations of DNA molecules carrying modifications like substitutions, small deletions, insertions or inversions, which

modifications like substitutions, small deletions, insertions or inversions, which nevertheless encode polypeptides having substantially the biochemical activity of the 29 kDa polypeptide. It will be known to the skilled person that  $A \leftrightarrow G$  and  $T \leftrightarrow C$  substitutions, with no effect on the amino acid sequence, are not unusual in H. pylori.

The only difference between SEQ ID NO: 1 and SEQ ID NO: 3 is that SEQ ID NO: 1 has an A residue in position 1458, while SEQ ID NO: 3 has a C residue in the same position.

Furthermore, the invention provides a formulation of DNA molecules which nucleotide sequences are degenerate, because of the genetic code, to the nucleotide sequence shown as SEQ ID NO: 1 or 3. Since there are 64 possible codons, but only 20 natural amino acids, most amino acids are coded for by more than one codon. This natural "degeneracy", or "redundancy", of the genetic code is well known in the art. It will thus be appreciated that the DNA sequence shown in the Sequence Listing is only an example within a large but definite group of DNA sequences which will encode the polypeptide as described above.

Consequently, the inventions includes also formulations based on a lipid aggregate of an isolated nucleic acid molecule selected from:

- (a) nucleic acid molecules comprising a nucleotide sequence which is identical with, or substantially similar to, positions 796-1572 or 874-1572 in SEQ ID NO: 1 or 3 in the Sequence Listing;
- (b) nucleic acid molecules comprising a nucleotide sequence capable of hybridizing to a nucleotide sequence complementary the polypeptide coding region of a DNA molecule as defined in (a) and which codes for a polypeptide as described above, or a functionally equivalent modified form thereof; and (c) nucleic acid molecules comprising a nucleic acid sequence which is degenerate as a result of the genetic code to a nucleotide sequence as defined in (a) or (b) and which codes for a polypeptide as described above, or a functionally equivalent modified form thereof.

### Brief description of the drawings

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Figure 1a) shows the result of eradication of *H. pylori* infection in mice after oral administration of different agents, such as a *Helicobacter* antigen alone or together with an adjuvant; the antigen is administered either directly (control) or incorporated in lipid aggregates according to the present invention. Legend to figure 1a) and abbrevations are as follow:

Number of *H. pylori* (as CFU, geometric mean values) in 25mm<sup>2</sup> scrapings of antrum and corpus, following oral administration of different agents.

CFU= Colony forming units; CT=Cholera toxin; LA=Lipid aggregates; LA(29 kDa)=29 kDa polypeptide formulated in Lipid aggregates.

Figure 1b) shows the same as Figure 1a) but displayed as the total number of *H. pylori* in the stomach. Abbrevations see: Figure 1a) above.

Figure 2) shows mucosal antibody response towards a *Helicobacter* antigen incorporated in a lipid aggregate according to the present invention. The figure 2) shows specifically the total amount of mucosal immunoglobulin (Ig) with a specificity for the antigen preparation

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comprising the 29 kDa polypeptide, detected by ELISA technique. Abbrevations see: Figure 1a) above.

Figure 3) shows mucosal antibody response towards a *Helicobacter* antigen incorporated in a lipid aggregate according to the present invention. The figure 3) shows specifically the amount of mucosal IgA antibodies with a specificity for the antigen preparation comprising the 29 kDa polypeptide. Abbrevations see: Figure 1a) above.

Figure 4) shows serum antibody response towards a *Helicobacter* antigen incorporated in a lipid aggregate according to the present invention. The figure 4) shows specifically the amount of Helicobacter specific IgG antibodies in sera with a specificity for an antigen preparation comprising a membrane protein. Abbrevations see: Figure 1a) above.

Figure 5) shows serum antibody response towards a *Helicobacter* antigen incorporated in a lipid aggregate according to the present invention. The figure 4) shows specifically the amount of 29 kDa specific IgG antibodies in sera with a specificity for an antigen preparation comprising 29 kDa polypeptide. Abbrevations see: Figure 1a) above.

#### Disclosure of the invention

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It has now been found that an improved immune response to *Helicobacter pylori* antigens as defined above can be obtained by administration of said antigens in a pharmaceutical formulation based on lipidic material, i.e. in the form of a lipid aggregate.

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Means of obtaining the improved formulations are based on the use of negatively charged lipids, either alone or in combination with other lipid material such as, but not limited to phospholipids, glycerides, etc, and in combination with a cation component, such as positively charged ions. The use of the formulations result in an improved immune response to the antigens described above when given by different

administration routes, such as, but not limited to, the oral, the rectal, the tonsillar, the buccal, the nasal, the vaginal etc, administration route. The preferred administrations are the peroral, rectal and nasal routes.

The negatively charged lipids can be, but are not limited to: phosphatidylserine, phosphatidylinositol, phosphatidic acid, and phosphatidylglycerol.

The said negatively charged lipids might be combined with one or more additives in the formulation, such as, but not limited to:

- other lipids, e g cholesterol, phosphatidylcholine, phosphatidyletanolamine, etc;
  - ions, e g Ca<sup>2+</sup>, Mg<sup>2+</sup>, Ba<sup>2+</sup>, Zn<sup>2+</sup>, Mn<sup>2+</sup>, Al<sup>3+</sup> etc.
  - adjuvants to further enhance the immunological response, e g lipid A and its derivatives, saponins like QS-21, cholera toxin (CT), etc.
- The process chosen for manufacturing of said pharmaceutical formulation is exemplified by the following general description, and in the Experimental Section below.

In general, these procedures are based on the formation of the lipid aggregates in three steps:

- the formation of a lipid film composed of the negatively charged lipids with or without other lipids, e.g. cholesterol;
- the addition of the antigen in a surfactant solution which then also produces mixed micelles with the lipids with incorporated antigen, preferably a nonionic surfactant with high critical micelle concentration is used;
  - the exchange of the surfactant with the positively charged ion.

The last step can be carried out in different ways. One way is to first remove the surfactant, e g by dialysis, and then add for example Ca<sup>2+</sup> in a second dialysis step, or

directly from a concentrated Ca<sup>2+</sup> solution. Another way is to exchange the surfactant with the positively charged ion in one dialysis step. A modification of these processes can be utilized if the antigen is insensitive to organic solvents. Thus, the antigen is dissolved together with the lipids in an organic solvent. The formation of liposomes with incorporated antigen is then formed after evaporation of the solvent and after the dispersion in a buffer solution. The lipid aggregates can finally be induced by the addition of positively charged ions by direct addition or by dialysis. Modifications of these processes are possible and can be chosen according to literature by those skilled in the art.

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The dosage form used may be a solid, semisolid, liquid dispersion, or solutions prepared by use of well known pharmaceutical techniques, such as blending, granulation, milling, spray drying, compaction, coating, etc. Further, the formulations may be monolithic, such as tablets, or capsules, or in the form of multiple formulations administered in a tablet, capsule or sachets.

The antigen is a polypeptide for use in therapy of *Helicobacter pylori* infection in a mammal, including man, and for use as a therapeutic or prophylactic vaccine as described above.

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Another important aspect of the invention is a vaccine formulation for inducing a protective immune response in a mammal, including humans, against the bacillary and/or coccoid form of *Helicobacter pylori*. Such a vaccine formulation comprises a lipid aggregate and an immunogenically effective amount of a polypeptide as defined above. Preferably the polypeptide includes at least a part of the 29 kDa polypeptide comprising an immunogenic epitope, or a modified form of said polypeptide which retains the capability to induce protective immunity against *Helicobacter pylori* infection. The term "modified form" includes, but is not restricted to, forms of the polypeptide which are post-translationally modified, e.g. lipidated. It is believed that the 29 kDa protein is lipidated.

The vaccine formulation according to the invention can be used for both therapeutic and prophylactic purposes. The vaccine formulation according to the invention is preferably administered to any mammalian mucosa exemplified by the buccal, the nasal, the tonsillar, the gastric, the intestinal (small and large intestine), the rectal and the vaginal mucosa. The mucosal vaccines can be given together with for the purpose appropriate adjuvants. The vaccine can also be given parenterally, by subcutaneous, intracutaneous or intramuscular route, optionally together with the appropriate adjuvant. With the term "adjuvant" is meant a component which in general enhances the effect of the vaccine. Such adjuvants are for instance cholera toxin and E.coli heatlabile toxin and their non-toxic derivatives saponins, cytokines and chemokines. Non-toxic derivatives of cholera toxin and E.coli heat labile toxin includes e.g. mutations with reduced ADP-ribosylating activity or CTA1-DD which binds only to  $1g^+$  cells.

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An alternative approach for creating an immune response against the 29 kDa polypeptide is to use the approach known as "nucleic acid vaccination" or "naked DNA" vaccination. It is known in the art that injection into muscle of plasmid DNA encoding an antigen of interest can result in sustained expression of the antigen and generation of an immune response (see e.g. Rabinovich et al. supra). Several routes of administration are possible, such as parental, mucosal or via a "gene-gun" that delivers tiny amounts of DNA-coated gold beads (Fynan et al. (1993) Proc. Natl. Acad. Sci. U.S.A. 90, 11478-11482).

Thus, a nucleic acid molecule can be expressed in plasmid comprising a suitable eukaryotic promoter. This "naked DNA" can then be given, e g orally with the formulation. Epitopes of the expressed protein will be expressed by MHC molecules on the surface of the cells and trigger an immune response. Consequently, nucleic acid molecules and vectors as disclosed in the previous paragraphs for use in therapy, in particular for use as a vaccine, are further aspects of the invention. The

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use of such nucleic acid molecules and vectors in the manufacture of compositions for treatment, prophylaxis or diagnosis of Helicobacter pylori infection are also further aspects of the invention.

In a further aspect, the invention provides a method of eliciting in a mammal, including man, a protective immune response against *Helicobacter pylori* infection, said method comprising the step of administering to the said mammal an immunologically effective amount of a vaccine formulation according to the present invention. The term "immunologically effective amount" is intended to mean an amount which elicit a significant protective *Helicobacter pylori* response, which will eradicate a *H. pylori* infection in an infected mammal or prevent the infection in a susceptible mammal. Typically an immunologically effective amount would comprise approximately 1 μg to 1 000 mg, preferably approximately 100 μg to 100 mg, of a *H. pylori* antigen for oral administration, or approximately less than 100 μg for parenteral administration. An immunologically effective amount of the antigen is typically administered in a formulation with the antigen in a range from 0.01 % w/w to 99 % w/w.

#### Detailed description of the invention

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The following examples illustrative different aspects of the invention. The preparation of the lipid aggregate comprising a *H. pylori* antigen is described in Example 1 and its use is illustrated in Example 2. Examples 3 to 7 describe the preparation and tests of a suitable antigen for the pharmaceutical formulation prepared in Example 1, i.e. an antigen for the formulation according to the present invention.

# Preparation and testing of a formulation according to the present invention

EXAMPLE 1: Formulation of the 29 kDa polypeptide rHpaA

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#### 1.1 Materials & Methods

Materials: BPS (brain phosphatidylserine, Avanti), CH (cholesterol), TES (Ntris(Hydroxymethyl)methyl-2-aminoethanesulfonic acid), L-histidine, NaCl, NOG (N-octyl-β-D-glucopyranoside), CaCl, 2H<sub>2</sub>O, NaOH, chloroform, and methanol were used as purchased. The water was ELGA quality (18.2 MΩ). The 29 kDa polypeptide and polyclonal rabbit-anti-29 kDa were obtained inhouse according to the examples described below.

Methods: In this specific example, the final composition of the formulation was intended to contain 1 mg/mL of the antigen, with a lipid/antigen weight ratio of 4/1, and with a total lipid concentration in the range of 2.5 - 3 mg/mL. Thus, a lipid film of BPS and CH (ratio 9/1 w/w) was obtained in a rotavapor system by evaporation of the chloroform/methanol (9/1 v/v) solvent that had been used for mixing the lipids. The lipid film was dissolved in a buffer solution (2 mM TES, 2 mM L-histidine, 100 mM NaCl, pH adjusted to 8.0 with NaOH(aq)) containing 2% w/w NOG (the NOG/lipid ratio intended to be 7.5/1 w/w) with or without the antigen. NOG was exchanged for Ca²+ during dialysis (Spectrapoor no 1, MW cutoff 6000-8000), twice versus a 3mM Ca²+ buffer solution (overnight and four hours), and twice versus a 6 mM Ca²+ solution (overnight and four hours).

After centrifugation of the contents, the supernatant and the pellet were analyzed for their antigen contents by different methods (Lowry assay, ELISA (polyclonal rabbitanti-29 kDa), SDS-PAGE, and Western blot). The following results were obtained:

- By Lowry: concentration in the total formulation: 0.41 ± 0.05 mg/mL; pellet: 0.41 ±0.04 mg/mL. In the supernatant as well as in formulations prepared without antigen the protein contents was below the level of detection;
  - By ELISA: the concentrations of the total formulation and the pellet were in the range of 0.6 1.2 mg/mL, while the supernatant concentration was 0.01-0.02

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mg/mL. In the formulations prepared without antigen the protein concentration was below the detection limit of the method;

- By SDS-PAGE: The 29 kDa polypeptide shows no significant change after formulation into lipid aggregates.
- By Western blot: Trace amounts of 29 kDa polypeptide were found in the supernatant of the formulation manufactured with antigen.

After analysis the concentration of the centrifugated formulation was adjusted with 6mM Ca<sup>2+</sup> buffer (vortex and 20 seconds in an ultrasonic bath) to give a composition in the formulation to be administered corresponding to 0.33 mg/mL of 29 kDa polypeptide. Administration of 300 µL would then give 100 µg administered 29 kDa polypeptide to each animal. Formulations manufactured without antigen were adjusted in the same way.

EXAMPLE 2: Analysis of the *H. pylori* 29 kDa protein for use as a vaccine, when used directly and when incorporated in a lipid aggregate formulation.

### 2.1. Materials & Methods

#### 20 2.1.1. Animals

Female SPF BALB/c mice were purchased from Bomholt Breeding centre (Denmark). They were kept in ordinary makrolon cages with free supply of water and food. The animals were 4-6 weeks old at arrival.

### 25 2.1.2. Infection

After a minimum of one week of acclimatization, the animals were infected with a type 2 strain of *H. pylori* (strain 244, originally isolated from an ulcer patient). This strain has earlier proven to be a good colonizer of the mouse stomach. The bacteria were grown overnight in Brucella broth supplemented with 10% fetal calf serum, at

+37°C in a microaerophilic atmosphere (10%  $CO_2$ , 5%  $O_2$ ). The animals were given an oral dose of omeprazole (400  $\mu$ mol/kg) and after 3-5 h an oral inoculation of H. pylori (approximately  $10^7$ - $10^8$  cfu/animal). Infection was checked in control animals 2-3 weeks after the inoculation.

2.1.3. Immunizations

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The 7 groups of mice, with 6 animals/group were immunized 4 times over a 34 day period (day 1, 15, 25 and 35). Recombinant purified Helicobacter 29 kDa polypeptide was given at a dose of 100  $\mu$ g/mouse either directly suspended in TES buffer or as incorporated into lipid aggregates (see above) . The groups are listed in Table I.

As an adjuvant, some groups of the animals were also given 10  $\mu$ g/mouse of cholera toxin (CT) with each immunization. A total volume of 0.3 ml was given at each immunization. Omeprazole (400  $\mu$ mol/kg) was given orally to the animals 3-5 h prior to immunization as a way of protecting the antigens from acid degradation. Animals were sacrificed 1-4 weeks after final immunization.

### 2.1.4. Analysis of infection

The mice were sacrificed by CO<sub>2</sub> and cervical dislocation. The abdomen was opened and the stomach removed. After cutting the stomach along the greater curvature, it was rinsed in saline. In in stomach half, an area of 25 mm² of the mucosa from the antrum and corpus was scraped separately with a surgical scalpel. The mucosa scraping was suspended in Brucella broth and plated onto Blood Skirrow plates. The plates were incubated under microaerophilic conditions for 3-5 days and the number of colonies was counted. The identity of *H. pylori* was ascertained by urease and catalase test and by direct microscopy or Gram staining.

#### 2.1.5 Analysis of Immune response

Mucosal antibodies were collected by the following technique. One half of the rinsed stomach was placed mucosal side up on a piece of paper. Likewise the duodenum was cut open and placed mucosal side up. One standardized round filter paper (30.4 mm²) was placed on the antrum and one on the corpus musosa. After 10 minutes the papers were transferred to one tube with 200µl special buffer containing protease inhibitors. A paperstrip 4.8x19 mm (91.2 mm²) was placed in the same way on the duodenum mucosa and subsequently treated in the same way. The buffer solution was, after a minimum of one hour extraction of the filter papers, used directly for ELISA measurements of antibody concentration or keep frozen at -20°C.

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<u>Serum antibodies</u> were collected from blood drawn by heart-puncture under - anesthesia. Prior to centrifugation, the blood was diluted with equal amount of PBS. The serum was kept at -20°C until analysis.

Mucosal antibodies were measured using an ELISA were 29 kDa polypeptide was plated followed by addition of mucosal extract. The response was developed by using Alkaline Phospahatase labelled anti-IgA and anti-Ig. Standad curves were created from plating known amounts of mouse IgA and Ig instead of the unknown sample. Serum antibodies were measured in a similar way but in this case both membrane proteins of the infecting strain i.e. 244 as well as 29 kDa polypeptide was plated.

#### 2.2. Results

#### 2.2.1. Eradication of infection

The animals in this study were infected with *H. pylori* strain 244 one month prior to immunizations. Mice in groups of six were then immunized with either cholera toxin (CT) or CT together with the *Helicobacter* 29 kDa polypeptide either directly or formulated in lipid aggregates (LA). Lipid aggregates were also tested without any

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protein incorporation with and without CT (see Table I) Control animals received vehicle only (TES buffer). One week after the final immunization, the animals were sacrificed and colony forming units (CFU) was determined (see Fig. 1a, and Fig. 1b). All control animals, as well as those receiving CT, LA and CT+LA were heavily infected in both antrum and corpus. Animals actively immunized with 29 kDa polypeptide plus CT, had significantly decreased bacterial content (CFU values) compared with the controls. In 2/5 mice no bacteria could be detected in the corpus. When the 29 kDa polypeptide was formulated in lipid aggregates according to the present invention the degree of *H. pylori* colonization in the gastric mucosa decreased compared to administration of lipid aggregate alone. The addition of CT to 29 kDa polypeptide containing lipid aggregate further decreased the bacterial content, especially in the antrum.

### 2.2.2 Mucosal antibody response

Specific mucosal antibodies towards 29 kDa polypeptide of general Ig and of IgA class was measured. In the general Ig class increased levels were seen following immunization with HpaA+CT, in the group were 29 kDa polypeptide was formulated with lipid aggregates and when CT was added to the latter preparation (see Figure 2). The appearence of specific IgA showed the same pattern (see Figure 3).

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### 2.2.3 Serum antibody response

Serum antibody response was measured towards a membrane prepartion of *H. pylori* strain 244 (the strain used for infection of the mice) and towards 29 kDa polypeptide. All animals had high titers against 244. No difference between the groups could be detected (see Figure 4). This response is mainly seen as a sign of appropriate infection with *H. pylori*. The specific serum response to 29 kDa polypeptide preparation as a sign of successful presentation of the 29 kDa polypeptide to the systemic immune system was highest in the group with 29 kDa + CT followed by 29 kDa polypeptide

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formulated with LA + CT. Also the lipid aggregate formulation of 29 kDa polypeptide without CT gave an increased number of serum antibodies (see Fig. 5).

### 2.3. Discussion and Applicability of the invention

It has previously been shown that the administration of 29 kDa + CT can give rise to an immune response measured as mucosal and systemic antibodies, which is capable of eradicate of decrease the number of *H. pylori* associated with the mouse gastric mucosa. In the absence of the adjuvant CT, no effect has been obtained.

Therefore, surprisingly it was found that 29 kDa polypeptide formulated in the lipid aggregates according to the present invention could, even in the absence of CT, induce an immune response of such nature that the degree of *H. pylori* colonization of the gastric mucosa of mice decreased.

#### 2.4. Conclusions

- The *H. pylori* surface located protein 29 kDa polypeptide has been shown to be a strong and consistent antigen when in a purified recombinant form is presented to a mucosal surface in a physiological buffer solution.
- The 29 kDa polypeptide will stimulate a competent local immune response capable of eradicating *H. pylori* colonizing the gastric mucosa.
  - Repeated administration of 29 kDa polypeptide during weeks or months will potentiate the effect.
  - The effect of 29 kDa polypeptide has been dependent on the simultaneous presence of the mucosal adjuvant Cholera toxin (CT).
  - Surprisingly 29 kDa polypeptide will induce an adequate local immune response even in the absence of CT, when it is formulated in lipid aggregates.

It is concluded that 29 kDa polypeptide can evoke a relevant immune response locally in the gastric mucosa both when administered together with cholera toxin as well as when it is formulated in a lipid aggregate.

This lipid aggregate formulation obviously has the capability to present *H. pylori* antigens to the local immune system in such a way that an adequate eradicating immune response is obtained.

TABLE I: Therapeutic immunisation groups

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Substance:	Dose:	Administra-	Number of		
		tion	doses		
1. Control, vehichle	0,3 ml/dose	p.o	4		
2. Cholera toxin (CT) 10µg	0,3 ml/dos	p.o	4		
3. Lipid aggregates (LA)	0,3 ml/dose	p.o	4		
4. LA + CT	0,3 ml/dose	p.o	4		
5. 29 kDa(100µg) + CT	0,3 ml/dose	p.o	4		
6. LA(29 kDa) 100 μg	0,3 ml/dose	p.o	4		
7. LA(29 kDa) + CT	0,3 ml/dose	p.o	4		

Further support is found in the accompaning figures as discussed above.

A suitable antigen for the formulation according to the present invention is the 29 kDa polypeptide described in the Applicant's co-pending WO96/38475. The 29 kDa polypeptide used in examples 1 and 2 was prepared according to the published WO96/38475.

#### **DEPOSITION OF MICROORGANISMS**

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The plasmid pAE1 has been deposited under the Budapest Treaty at the National Collections of Industrial and Marine Bacteria (NCIMB), Aberdeen, Scotland, UK, and under accession number NCIMB 40732. The date of deposit is 16 May 1995.

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#### SEQUENCE LISTING

(1) GENERAL INFORMATION:

5

- (i) APPLICANT:
  - (A) NAME: ASTRA AB
  - (B) STREET: Västra Mälarehamnen 9
  - (C) CITY: Södertälje

10

- (E) COUNTRY: Sweden
- (F) POSTAL CODE (ZIP): S-151 85
- (G) TELEPHONE: +46 8 553 260 00
- (H) TELEFAX: +46 8 553 288 20
- (I) TELEX: 19237 astra s

15

- (ii) TITLE OF INVENTION: New Pharmaceutical Formulation of Polypeptides
- (iii) NUMBER OF SEQUENCES: 4
- 20 (iv) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: Floppy disk
  - (B) COMPUTER: IBM PC compatible
  - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
  - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)

25

(2) INFORMATION FOR SEQ ID NO: 1:

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(i) SEQUENCE CHARACTERISTICS:

	(A) LENGTH: 1670 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: double	
5	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: cDNA	
10	(ix) FEATURE:	
	(A) NAME/KEY: CDS	
	(B) LOCATION:7931575	
	(ix) FEATURE:	
15	(A) NAME/KEY: mat_peptide	
	(B) LOCATION: 7931572	
	/ ') anaumyan nuganyanyan ana in wa i	
••	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:	
20	GATCCTATCG CGCCAAAGGT GGTATTAGGA ATAAGAGCTT GATTATTAAT CTCCCTGGTA	60
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	AGTCCAAAAA GTATTAGAGA ATGCTTAGAG GCGGTTTTTC CAGCGATTCC TTATTGCGTG	120
25	GATTTGATTT TAGGGAATTA CATGCAAGTG AATGAAAAAA ACATTCAAGC GTTTGCCCCC	180
	AAACAATAAG GTAAAAAATG CCACTCACTC ATTTGAATGA AGAAAATCAA CCTAAAATGG	240

•	TGGATATAGG	GGATAAAGAA	ACCACTGAAA	GAATCGCTCT	AGCAAGCGGT	CGTATCAGCA	300
	TGAATAAAGA	GGCTTATGAC	GCTATTATCA	ATCATGGCGT	CAAAAAGGGT	CCGGTATTAC	360
5	AAACTGCTAT	TATTGCTGGG	ATTATGGGGG	CTAAAAAGAC	AAGCGAACTC	ATTCCCATGT	420
	GCCATCCAAT	CATGCTCAAT	GGGGTGGATA	TTGATATTTT	AGAAGAAAA	GAGACTTGTA	480
	GTTTTAAACT	CTATGCGAGA	GTCAAAACTC	AAGCTAAAAC	GGGCGTAGAA	ATGGAAGCGC	540
10	TAATGAGTGT	GAGCGTAGGG	CTTTTAACCA	TTTATGACAT	GGTGAAAGCC	ATTGATAAGA	600
	GCATGACAAT	TAGCGGTGTG	ATGCTGGAAT	ATAAAAGTGG	AGGCAAAAGT	GGGGATTATA	660
15	ACGCTAAAAA	ATAGAAAAAG	ACTGATAATC	TAAAGATATT	AGGGTAAAAT	AACATTTTGA	720
	CAACAAAAGC	GTGTTGGTTG	CTTCGGATTT	GTTGTTATAG	AAGTCTAAAA	TATTACAATC	780
	AAGGATAGAA						828
20		Met Arg	Ala Ash Ash 5		rs Asp Phe A	la Trp	
	AAA AAA TGC	CTT TTA GG	c gcg agc g	TG GTG GCT	TTA TTA GTG	GGA TGC	876
25	Lys Lys Cys		y Ala Ser V 20	al Val Ala		Gly Cys	
25	15		20		25		
	AGC CCG CAT						924
	Ser Pro His	Ile Ile Gl	u Thr Asn G	lu Val Ala	Leu Lys Leu	Asn Tvr	

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		30					35					40					
	CAT	CCA	GCT	AGC	GAG	AAA	GTT	CAA	GCG	TTA	GAT	GAA	AAG	ATT	TTG	CTT	972
	His	Pro	Ala	Ser	Glu	Lys	Val	Gln	Ala	Leu	Asp	Glu	Lys	Ile	Leu	Leu	
5	45					50					55					60	
	тта	AGG	CCA	GCT	ттс	CAA	тат	AGC	GAT	аат	ልጥ <b>ሮ</b>	CCT	מממ	GAG	ጥልጥ	GAA	1020
				Ala													, 1020
	Deu	n.y	110	ALU	65	GI.I.	131	Der	ಬಾಗಿ	70	TIC	AIG	шys	GIU		GIU	
10					03					70					75		
10						~											•
				AAG													1068
	Asn	Lys	Phe	Lys	Asn	Gln	Thr	Ala	Leu	Lys	Val	Glu	Gln	Ile	Leu	Gln	
				80					85					90			
15	AAT	CAA	GGC	TAT	AAG	GTT	ATT	AGC	GTA	GAT	AGC	AGC	GAT	AAA	GAC	GAT	1116
	Asn	Gln	Gly	Tyr	Lys	Val	Ile	Ser	Val	Asp	Ser	Ser	Asp	Lys	Asp	Asp	
		•	95					100					105				
	TTT	TCT	TTT	GCA	CAA	AAA	AAA	GAA	GGG	TAT	TTG	GCG	GTT	GCT	ATG	AAT	1164
20	Phe	Ser	Phe	Ala	Gln	Lys	Lys	Glu	Gly	Tyr	Leu	Ala	Val	Ala	Met	Asn	
		110					115					120					
	GGC	GAA	ATT	GTT	TTA	CGC	CCC	GAT	CCT	AAA	AGG	ACC	ATA	CAG	AAA	AAA	1212
	Gly	Glu	Ile	Val	Leu	Arg	Pro	Asp	Pro	Lys	Arg	Thr	Ile	Gln	Lys	Lvs	
25	125					130					135				-	140	
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	тса	GD A	ccc	GGG	מיזית	ጥጥል	ጥጥ	ጥርር	»CC	രത	ጥጥር፤	CAC	**	אתיי	C	GGG	1050
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Ser Glu Pro Gly Leu Leu Phe Ser Thr Gly Leu Asp Lys Met Glu Gly

GTT TTA ATC CCG GCT GGG TTT ATT AAG GTT ACC ATA CTA GAG CCT ATG Val Leu Ile Pro Ala Gly Phe Ile Lys Val Thr Ile Leu Glu Pro Met AGT GGG GAA TCT TTG GAT TCT TTT ACG ATG GAT TTG AGC GAG TTG GAC Ser Gly Glu Ser Leu Asp Ser Phe Thr Met Asp Leu Ser Glu Leu Asp ATT CAA GAA AAA TTC TTA AAA ACC ACC CAT TCA AGC CAT AGC GGG GGG Ile Gln Glu Lys Phe Leu Lys Thr Thr His Ser Ser His Ser Gly Gly TTA GTT AGC ACT ATG GTT AAG GGA ACG GAT AAT TCT AAT GAC GCG ATC ·\*15 Leu Val Ser Thr Met Val Lys Gly Thr Asp Asn Ser Asn Asp Ala Ile AAG AGC GCT TTG AAT AAG ATT TTT GCA AAT ATC ATG CAA GAA ATA GAC Lys Ser Ala Leu Asn Lys Ile Phe Ala Asn Ile Met Gln Glu Ile Asp AAA AAA CTC ACT CAA AAG AAT TTA GAA TCT TAT CAA AAA GAC GCC AAA Lys Lys Leu Thr Gln Lys Asn Leu Glu Ser Tyr Gln Lys Asp Ala Lys 

GAA TTA AAA GGC AAA AGA AAC CGA TAA AAACAAATAA CGCATAAGAA

Glu Leu Lys Gly Lys Arg Asn Arg \*

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255 260

AAGAACGCTT GAATAAACTG CTTAAAAAGG GTTTTTTAGC GTTCTTTTTG AGCGTGTATT 1655

5 TAAGGGCTGA TGATC 1670

(2) INFORMATION FOR SEQ ID NO: 2:

10 (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 261 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

15 (ii) MOLECULE TYPE: protein

20

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

Met Arg Ala Asn Asn His Phe Lys Asp Phe Ala Trp Lys Lys Cys Leu

1 5 10 15

Leu Gly Ala Ser Val Val Ala Leu Leu Val Gly Cys Ser Pro His Ile

20 25 30

Ile Glu Thr Asn Glu Val Ala Leu Lys Leu Asn Tyr His Pro Ala Ser

25 35 40 45

Glu Lys Val Gln Ala Leu Asp Glu Lys Ile Leu Leu Leu Arg Pro Ala

50 55 60

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Phe	Gln	Tyr	Ser	Asp	Asn	Ile	Ala	Lys	Glu	Tyr	Glu	Asn	Lys	Phe	Lys
65					70					75					80
Asn	Gln	Thr	Ala	Leu 85	Lys	Val	Glu	Gln	Ile 90	Leu	Gln	Asn	Gln	Gly 95	-
Lys	Val	Ile	Ser	Val	Asp	Ser	Ser		Lys	Asp	Asp	Phe		Phe	Ala
-	_	_		4.1		_									
Gln	Lys	Lys 115	Glu	GIĀ	Tyr	Leu	120	Val	Ala	Met	Asn	Gly 125	Glu	Ile	Val
Leu		Pro	Asp	Pro	Lys		Thr	Ile	Gln	Lys		Ser	Glu	Pro	Gly
	130					135					140				
	Leu	Phe	Ser	Thr		Leu	Asp	Lys	Met		Gly	Val	Leu	Ile	Pro 160
															100
Ala	Gly	Phe	Ile	Lys 165	Val	Thr	Ile	Leu	Glu 170	Pro	Met	Ser	Gly	Glu 175	Ser
Leu	Asp	Ser	Phe	Thr	Met	Asp	Leu	Ser 185	Glu	Leu	Asp	Ile	Gln 190	Glu	Lys
Phe	Leu	Lys 195	Thr	Thr	His		Ser 200	His	Ser	Gly			Val	Ser	Thr
	Asn Lys Gln Leu 145 Ala	Asn Gln Lys Val Leu Arg 130 Leu Leu 145 Ala Gly Leu Asp	Asn Gln Thr Lys Val Ile Gln Lys Lys 115 Leu Arg Pro 130 Leu Leu Phe 145 Ala Gly Phe Leu Asp Ser	Asn Gln Thr Ala Lys Val Ile Ser 100 Gln Lys Lys Glu 115 Leu Arg Pro Asp 130 Leu Leu Phe Ser 145 Ala Gly Phe Ile Leu Asp Ser Phe 180 Phe Leu Lys Thr	Asn Gln Thr Ala Leu 85 Lys Val Ile Ser Val 100 Gln Lys Lys Glu Gly 115 Leu Arg Pro Asp Pro 130 Leu Leu Phe Ser Thr 145 Ala Gly Phe Ile Lys 165 Leu Asp Ser Phe Thr 180 Phe Leu Lys Thr Thr	Asn Gln Thr Ala Leu Lys 85         Lys Val Ile Ser Val Asp 100         Gln Lys Lys Glu Gly Tyr 115         Leu Arg Pro Asp Pro Lys 130         Leu Leu Phe Ser Thr Gly 150         Ala Gly Phe Ile Lys Val 165         Leu Asp Ser Phe Thr Met 180         Phe Leu Lys Thr Thr His	Asn Gln Thr Ala Leu Lys Valast         Lys Val Ile Ser Val Asp Ser 100         Gln Lys Lys Glu Gly Tyr Leu 115         Leu Arg Pro Asp Pro Lys Arg 130         Leu Leu Phe Ser Thr Gly Leu 145         Ala Gly Phe Ile Lys Val Thr 165         Leu Asp Ser Phe Thr Met Asp 180         Phe Leu Lys Thr Thr His Ser	Asn Gln       Thr Ala Leu Lys Val Glu 85         Lys Val       Ile Ser Val Asp Ser Ser 100         Gln       Lys Lys Glu Gly Tyr Leu Ala 115         Leu Arg Pro Asp Pro Lys Arg Thr 130         Leu Leu Phe Ser Thr Gly Leu Asp 145         145       150         Ala Gly Phe Ile Lys Val Thr Ile 165         Leu Asp Ser Phe Thr Met Asp Leu 180         Phe Leu Lys Thr Thr His Ser Ser	Asn       Gln       Thr       Ala       Leu       Lys       Val       Glu       Gln         Lys       Val       Ile       Ser       Val       Asp       Ser       Ser       Asp         Lys       Val       Ile       Ser       Val       Asp       Ser       Ser       Asp         Gln       Lys       Lys       Glu       Gly       Tyr       Leu       Ala       Val         Leu       Arg       Pro       Asp       Pro       Lys       Arg       Thr       Ile         Leu       Arg       Pro       Asp       Pro       Lys       Arg       Thr       Ile         Leu       Leu       Phe       Ser       Thr       Gly       Leu       Asp       Lys         Ala       Gly       Phe       Ile       Lys       Val       Thr       Ile       Leu         Leu       Asp       Ser       Phe       Thr       Met       Asp       Leu       Ser         Leu       Asp       Ser       Phe       Thr       Met       Asp       Leu       Ser         Leu       Asp       Ser       Phe       Thr       Met       A	65       The Ala Leu Lys Val Glu Gln Ile 85         Asn Gln Thr Ala Leu Lys Val Glu Gln Ile 85       85       Val Glu Gln Ile 90         Lys Val Ile Ser Val Asp Ser Ser Asp Lys 105       105       105         Gln Lys Lys Glu Glu Gly Tyr Leu Ala Val Ala 115       105       105         Leu Arg Pro Asp Pro Lys Arg Thr Ile Gln 130       135       120         Leu Leu Phe Ser Thr Gly Leu Asp Lys Met 145       150       150         Ala Gly Phe Ile Lys Val Thr Ibr Leu Glu 165       170         Leu Asp Ser Phe Thr Met Asp Leu Ser Glu 180       185         Phe Leu Lys Thr Thr His Ser Ser His Ser	Asn Gln       Thr       Ala       Leu Lys       Val       Glu       Gln       Ile       Leu Leu Rys       Ser       Ser       Asp       10e       Leu Leu Rys       Asp       Ser       Asp       Lys       Asp       Asp	Asn Gln       Thr       Ala       Leu       Lys       Val       Glu       Gln       Ile       Leu       Gln         Lys       Val       Ile       Ser       Val       Asp       Ser       Asp       Lys       Asp       Asp         Lys       Val       Ile       Ser       Val       Asp       Ser       Asp       Lys       Asp       Asp         Gln       Lys       Glu       Gly       Tyr       Leu       Ala       Val       Ala       Met       Asp       Asp         Leu       Arg       Pro       Asp       Pro       Lys       Arg       Thr       Ile       Gln       Lys       Lys         Leu       Arg       Pro       Asp       Pro       Lys       Arg       Thr       Ile       Gln       Lys       Lys         Ala       Gly       Pro       Pro       Ins       Val       Thr       Ile       Leu       Arg       Ins       Ins       Ins         Leu       Arg       Pro       Ins       <	Asn Gln Thr       Ala Leu Lys Val Glu Gln Ile Leu Gln Asn 85 90         Lys Val Ile Ser Val Asp Ser Ser Asp Lys Asp Asp Phe 100       105         Gln Lys Lys Glu Gly Tyr Leu Ala Val Ala Met Asn Gly 115       120         Leu Arg Pro Asp Pro Lys Arg Thr Ile Gln Lys Lys Ser 130       135         Leu Leu Phe Ser Thr Gly Leu Asp Lys Met Glu Gly Val 145       150         Ala Gly Phe Ile Lys Val Thr Ile Leu Glu Pro Met Ser 165       170         Leu Asp Ser Phe Thr Met Asp Leu Ser Glu Leu Asp Ile 180       185         Phe Leu Lys Thr Thr His Ser Ser His Ser Gly Gly Leu	Asn Gln Thr       Ala Leu Lys Val Glu Gln Ile Leu Gln Asn Gln 85         Lys Val Ile Ser Val Asp Ser Gr Asp Lys Asp Asp Phe Ser 100         Gln Lys Lys Glu Gly Tyr Leu Ala Val Ala Met Asn Gly Glu 115         Leu Arg Pro Asp Pro Lys Arg Thr Ile Gln Lys Lys Ser Glu 130         Leu Leu Phe Ser Thr Gly Leu Asp Lys Met Glu Gly Val Leu 145         Ala Gly Phe Ile Lys Val Thr Ise Leu Glu Pro Met Ser Gly 165         Ala Gly Phe Ile Lys Val Thr Ile Leu Glu Pro Met Ser Gly 165         Leu Asp Ser Phe Thr Met Asp Leu Ser Glu Leu Asp Ile Gln 190         Phe Leu Lys Thr Thr His Ser Ser His Ser Gly Gly Leu Val	Asn Gln Thr Ala Leu Lys Val Glu Gln Ile Leu Gln Asn Gln Gly 85 90 95  Lys Val Ile Ser Val Asp Ser Ser Asp Lys Asp Asp Asp Phe Ser Phe 100 105 110  Gln Lys Lys Glu Gly Tyr Leu Ala Val Ala Met Asn Gly Glu Ile 115 120 125  Leu Arg Pro Asp Pro Lys Arg Thr Ile Gln Lys Lys Ser Glu Pro 130 135 140  Leu Leu Phe Ser Thr Gly Leu Asp Lys Met Glu Gly Val Leu Ile 145 150 155  Ala Gly Phe Ile Lys Val Thr Ile Leu Glu Pro Met Ser Gly Glu 165 170 175  Leu Asp Ser Phe Thr Met Asp Leu Ser Glu Leu Asp Ile Gln Glu 180 185 190  Phe Leu Lys Thr Thr His Ser Ser His Ser Gly Gly Leu Val Ser

Met Val Lys Gly Thr Asp Asn Ser Asn Asp Ala Ile Lys Ser Ala Leu
210 215 220

Asn Lys Ile Phe Ala Asn Ile Met Gln Glu Ile Asp Lys Lys Leu Thr
5 225 230 235 240

Gln Lys Asn Leu Glu Ser Tyr Gln Lys Asp Ala Lys Glu Leu Lys Gly

245 250 255

10 Lys Arg Asn Arg \*

260

- (2) INFORMATION FOR SEQ ID NO: 3:
- (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 1670 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: double

(D) TOPOLOGY: linear

20

- (ii) MOLECULE TYPE: cDNA
- (ix) FEATURE:

25 (A) NAME/KEY: CDS

(B) LOCATION: 793..1575

(ix) FEATURE:

(A) NAME/KEY: mat\_peptide

(B) LOCATION: 793..1572

### 5 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

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10	GATTTGATTT	TAGGGAATTA	CATGCAAGTG	AATGAAAAA	ACATTCAAGC	GTTTGCCCCC	180
	AAACAATAAG	GTAAAAAATG	CCACTCACTC	ATTTGAATGA	AGAAAATCAA	CCTAAAATGG	240
15	TGGATATAGG	GGATAAAGAA	ACCACTGAAA	GAATCGCTCT	AGCAAGCGGT	CGTATCAGCA	300
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	AAACTGCTAT	TATTGCTGGG	ATTATGGGGG	CTAAAAAGAC	AAGCGAACTC	ATTCCCATGT	420
20	GCCATCCAAT	CATGCTCAAT	GGGGTGGATA	TTGATATTTT	AGAAGAAAA	GAGACTTGTA	480
	GTTTTAAACT	CTATGCGAGA	GTCAAAACTC	AAGCTAAAAC	GGGCGTAGAA	ATGGAAGCGC	540
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	GCATGACAAT	TAGCGGTGTG	ATGCTGGAAT	ATAAAAGTGG	AGGCAAAAGT	GGGGATTATA	660

	ACG	CTAA	AAA A	ATAG	AAAA	AG A	CTGA'	TAAT(	C TA	AAGA'	TATT	AGG	GTAA.	AAT .	AACA'	TTTT	GA	720
	CAAC	CAAA	AGC (	GTGT'	rggt'	rg C	rtcg	GATT'	r gt	rgtt	ATAG	AAG'	rcta.	AAA '	TATT.	ACAA!	rc	780
5	AAGO	GATA	GAA (	CG A														828
				Me	et A:	rg A	la A	sn A	sn H	is P	he L	ys A:	sp P	he A	la T	rp		
					1				5					10				
				CTT														876
10	Lys	Lys	Cys	Leu	Leu	Gly	Ala	Ser	Val	Val	Ala	Leu	Leu	Val	Gly	Суз		
			15					20					25					
	AGC	CCG	CAT	ATT	ATT	GAA	ACC	AAT	GAA	GTC	GCT	TTG	AAA	TTG	AAT	TAC		924
	Ser	Pro	His	Ile	Ile	Glu	Thr	Asn	Glu	Val	Ala	Leu	Lys	Leu	Asn	Tyr		
45		30					35					40						
	CAT	CCA	GCT	AGC	GAG	AAA	GTT	CAA	GCG	TTA	GAT	GAA	AAG	ATT	TTG	CTT		972
	His	Pro	Ala	Ser	Glu	Lys	Val	Gln	Ala	Leu	Asp	Glu	Lys	Ile	Leu	Leu		
	45					50					55					60		
20																		
	TTA	AGG	CCA	GCT	TTC	CAA	TAT	AGC	GAT	AAT	ATC	GCT	AAA	GAG	TAT	GAA	:	1020
	Leu	Arg	Pro	Ala	Phe	Gln	Tyr	Ser	Asp	Asn	Ile	Ala	Lys	Glu	Tyr	Glu		
					65					70					75			
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	Asn	Lys	Phe	Lys	Asn	Gln	Thr	Ala	Leu	Lys	Val	Glu	Gln	Ile	Leu	Gln		
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	AAT	CAA	GGC	TAT	AAG	GTT	ATT	AGC	GTA	GAT	AGC	AGC	GAT	AAA	GAC	GAT	1116
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5	TTT	TCT	TTT	GCA	CAA	AAA	AAA	GAA	GGG	TAT	TTG	GCG	GTT	GCT	ATG	AAT	1164
	Phe	Ser	Phe	Ala	Gln	Lys	Lys	Glu	Gly	Tyr	Leu	Ala	Val	Ala	Met	Asn	
		110					115					120				•	
	GGC	GAA	ATT	GTT	TTA	CGC	CCC	GAT	CCT	AAA	AGG	ACC	ATA	CAG	AAA	AAA	1212
10	Gly	Glu	Ile	Val	Leu	Arg	Pro	Asp	Pro	Lys	Arg	Thr	Ile	Gln	Lys	Lys	
	125					130					135					140	
																GGG	1260
	Ser	Glu	Pro	Gly	Leu	Leu	Phe	Ser	Thr	Gly	Leu	Asp	Lys	Met	Glu	Gly	
15					145					150					155		
	GTT	TTA	ATC	CCG	GCT	GGG	TTT	ATT	AAG	GTT	ACC	ATA	CTA	GAG	CCT	ATG	1308
	Val	Leu	Ile		Ala	Gly	Phe	Ile		Val	Thr	Ile	Leu	Glu	Pro	Met	
				160					165					170			
20																	
													AGC				1356
	Ser	Gly		Ser	Leu	Asp	Ser		Thr	Met	Asp	Leu	Ser	Glu	Leu	Asp	
			175					180					185				
•-										<b>-</b>							
25													CAT				1404
	Ile		Glu	ŗàs	Phe	Leu		Thr	Thr	His	Ser		His	Ser	Gly	Gly	
		190					195					200					

	TTA	GTT	AGC	ACT	ATG	GTT	AAG	GGA	ACG	GAT	AAT	TCT	AAT	GAC	GCG	ATC	1452
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	205					210					215					220	
5	AAG	AGA	GCT	TTG	AAT	AAG	ATT	TTT	GCA	AAT	ATC	ATG	CAA	GAA	ATA	GAC	1500
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					225					230					235		
	AAA	AAA	CTC	ACT	CAA	AAG	ААТ	TTA	GAA	тст	TAT	CAA	AAA	GAC	GCC	AAA	1548
10	Lys	Lys	Leu	Thr	Gln	Lys	Asn	Leu	Glu	Ser	Tyr	Gln	Lys	Asp	Ala	Lys	
				240					245					250			
	GAA	TTA	AAA	GGC	AAA	AGA	AAC	CGA	TAA	AAAC	CAAAT	TAA C	GCAT	TAAGA	<b>L</b> A		1595
	Glu	Leu	Lys	Gly	Lys	Arg	Asn	Arg	*								
15			255					260									
	AAG	AACG	CTT (	SAATA	LAAC1	G CI	TAAT	AAAGO	GTT	rTTT	ragc	GTTC	TTT	rtg <i>e</i>	\GCG1	GTATT	1655
	TAAC	GGC'	rga :	rgato	2												1670
20																	
	(2)	INFO	ORMA	rion	FOR	SEQ	ID N	IO: 4	l:								
	•				•	-											
			(i) 9	SEQUE	ENCE	CHAR	LACTE	ERIST	ICS:								
25				A) LE													
			1.5	.,		20											

(B) TYPE: amino acid

(D) TOPOLOGY: linear

							prot IPTI		SEQ :	ID NO	D: 4	:				
5	Met 1	Arg	Ala	Asn	Asn 5	His	Phe	Lys	Asp	Phe	Ala	Trp	Lys	Lys	Cys 15	Leu
	Leu	Gly	Ala	Ser 20	Val	Val	Ala	Leu	Leu 25	Val	Gly	Суз	Ser	Pro 30	His	Ile
10	Ile	Glu	Thr	Asn	Glu	Val	Ala	Leu 40	Lys	Leu	Asn	Tyr	His	Pro	Ala	Ser
	Glu	Lys 50	Val	Gln	Ala	Leu	Asp 55	Glu	Lys	Ile	Leu	Leu 60	Leu	Arg	Pro	Ala
15	Phe	Gln	Tyr	Ser	Asp	Asn 70	Ile	Ala	Lys	Glu	Tyr 75	Glu	Asn	Lys	Phe	Eys
20	Asn	Gln	Thr	Ala	Leu 85	Lys	Val	Glu	Gln	Ile 90	Leu	Gln	Asn	Gln	Gly 95	Tyr
	Lys	Val	Ile	Ser	Val	Asp	Ser	Ser	Asp 105	Lys	Asp	Asp	Phe	Ser	Phe	Ala

Leu Arg Pro Asp Pro Lys Arg Thr Ile Gln Lys Lys Ser Glu Pro Gly

125

25 Gln Lys Lys Glu Gly Tyr Leu Ala Val Ala Met Asn Gly Glu Ile Val

120

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130

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Leu Leu Phe Ser Thr Gly Leu Asp Lys Met Glu Gly Val Leu Ile Pro

145 150 155 160

Ala Gly Phe Ile Lys Val Thr Ile Leu Glu Pro Met Ser Gly Glu Ser

165 170 175

Leu Asp Ser Phe Thr Met Asp Leu Ser Glu Leu Asp Ile Gln Glu Lys

10 180 185 190

Phe Leu Lys Thr Thr His Ser Ser His Ser Gly Gly Leu Val Ser Thr

195 200 205

Met Val Lys Gly Thr Asp Asn Ser Asn Asp Ala Ile Lys Arg Ala Leu

210 215 220

Asn Lys Ile Phe Ala Asn Ile Met Gln Glu Ile Asp Lys Lys Leu Thr

225 230 235 240

Gln Lys Asn Leu Glu Ser Tyr Gln Lys Asp Ala Lys Glu Leu Lys Gly

245 250 255

Lys Arg Asn Arg \*

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## Legends to the figures:

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Fig.1a: Number of *Heliocobacter pylori* (as CFU, geometric mean values) in 25mm<sup>2</sup> scrapings of antrum and corpus, following oral administration of different agents.

CT=Cholera toxin; LA=Lipid aggregates; LA(29kDa)=29 kDa formulated in Lipid aggregates. n=6 for all animals, except CT (n=3) and 29kDa + CT (n=5).

Fig.1b: As 1a) but displayes as total number of *Heliocobacter pylori* in the stomach. Abbrevations see: Fig.1a)

Fig.2: Total amount of mucosal immunoglobulin (Ig) with a specificity for 29kDa polypeptide, detected by ELISA technique. Abbrevations see: Fig.1a

Fig.3: Amount of 29kDa specific mucosal IgA antibodies. Abbrevations see: Fig.1a

Fig.4: Heliocobacter specific IgG antibodies in sera. Abbrevations see: Fig.1a

Fig.5: 29kDa specific IgG antibodies in sera. Abbrevations see: Fig.1a

#### **CLAIMS**

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- 1. A pharmaceutical formulation comprising a lipid aggregate of a negatively charged lipid component or a lipid mixture and at least one antigenic polypeptide which is a *Helicobacter pylori* antigen or nucleic acid molecules coding for such polypeptides, and optionally a pharmaceutically acceptable carrier.
- 2. A pharmaceutical formulation according to claim 1, wherein the antigenic polypeptide is a recombinant polypeptide.
- 3. A pharmaceutical formulation according to claim 2, wherein the recombinant polypeptide is a hydrophobic *H. pylori* antigen.
  - 4. A pharmaceutical formulation according to claim 3, wherein the antigen is a *H. pylori* membrane protein.
  - 5. A pharmaceutical formulation according to claim 4, wherein the antigen is a polypeptide which has an amino acid sequence identical with, or substantially similar to, a *H. pylori* surface-exposed antigen with an approximate molecular weight of 29 kDa.
  - 6. A pharmaceutical formulation according to claim 5, wherein the polypeptide has an amino acid sequence identical with, or substantially similar to, SEQ ID NO:2 in the Sequence Listing.
- 7. A pharmaceutical formulation according to claim 1, wherein the polypeptide is a peptide with a length of at least 5 amino acids comprising an immunogenic epitope of a polypeptide which is a Helicobacter pylori membrane protein.

- 8. A pharmaceutical formulation according to claim 1, wherein the polypeptide is replaced by an isolated and purified nucleic acid molecule which has a nucleotide sequence coding for a polypeptide according to claim 5 or claim 6.
- 9. A pharmaceutical formulation according to claim 1, wherein the lipid aggregate in addition to the antigenic *H. pylori* polypeptide comprises a negatively charged lipid component or a lipid mixture comprises a cation component.
- 10. A pharmaceutical formulation according to claim 9, wherein the negatively charged lipid component is a phospholipid.
  - 11. A pharmaceutical formulation according to claim 10, wherein the phospholipid is selected from the group consiting of phosphatidylserine, phosphatidylinositol, phosphatidic acid and phosphatidylglycerol.
  - 12. A pharmaceutical formulation according to any of claims 9, 10 or 11, wherein there is also present a cation component which chelates or complexes with the negatively charged lipid component.
- <sup>20</sup> 13. A pharmaceutical formulation according to claim 12, wherein the cation component is selected from the group consisting of Ca<sup>2+</sup>, Mg<sup>2+</sup>, Ba<sup>2+</sup>, Zn<sup>2+</sup>, Mg<sup>2+</sup> and Al<sup>3+</sup>.
- 14. A pharmaceutical formulation according to claim 1, wherein the formulation in addition to the lipid aggregate comprises a vaccine adjuvant.
  - 15. A pharmaceutical formulation according to claim 1, for use as a therapeutic vaccine in a mammal, including man, which is infected by *Helicobacter pylori*.

- 16. A pharmaceutical formulation according to claim 1, for use as a prophylactic vaccine in a mammal, including man, from infection by *Helicobacter pylori*.
- 17. A pharmaceutical formulation according to claim 1, wherein the formulation is delivered to a mucosal surface by oral, parental, rectal or nasal administration.
  - 18. A process for the manufacture of a pharmaceutical formulation comprising a lipid aggregate of a negatively charged lipid component or a lipid mixture and at least one antigenic polypeptide which is *H. pylori* antigen, comprising the steps of
    - i) formation of a lipid film composed of the negatively charged lipid component or lipid mixture, optionally with other lipids,
    - ii) addition of the H. pylori antigen in a surfactant solution
    - iii) exchange the surfactant with a positively charged ion.
  - 19. A process according to claim 18, wherein step iii) is carried out by dialysis in one step, or in two steps.
- 20. Use of a formulation according to anyone of claims 1 to 18 in the manufacture of a formulation for the treatment or protection of *H. pylori* infections.
  - 21. Use of a formulation according to anyone of claims 1 to 18 in the manufacture of a vaccine eliciting a protective immune response against *H. pylori* infections.
- 25 22. A method of eliciting in a mammal a protective immune response against H. pylori infections, comprising the step of administering to the said mammal an immunogenically effective amount of a pharmaceutical formulation according to any of claims 1 to 18.

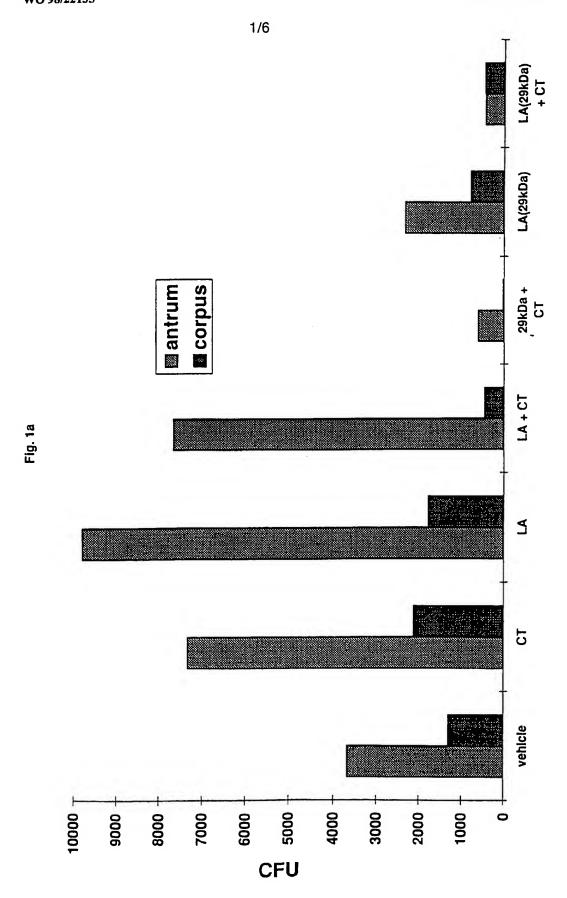
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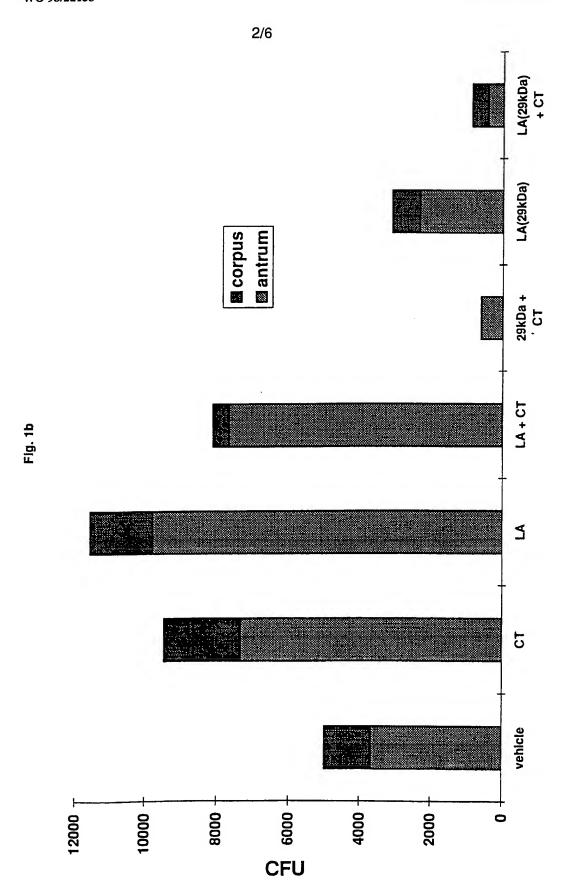
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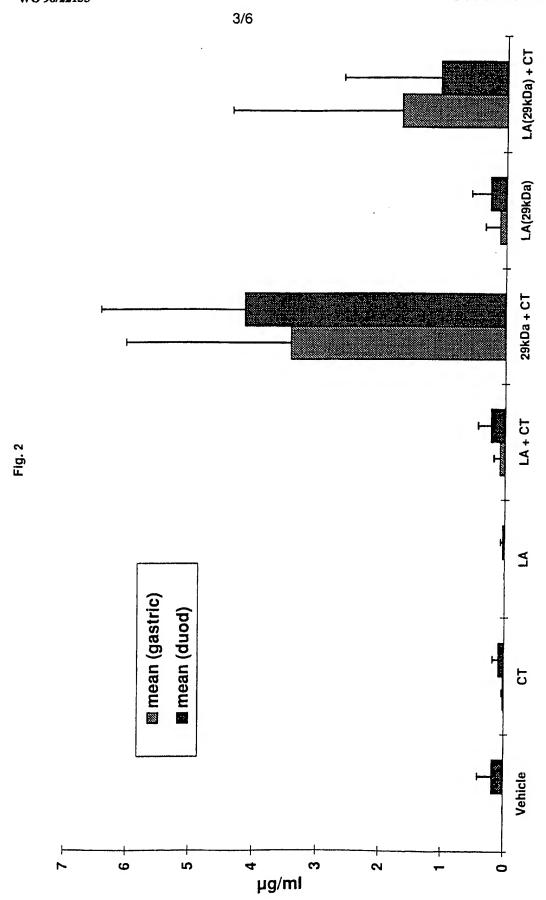
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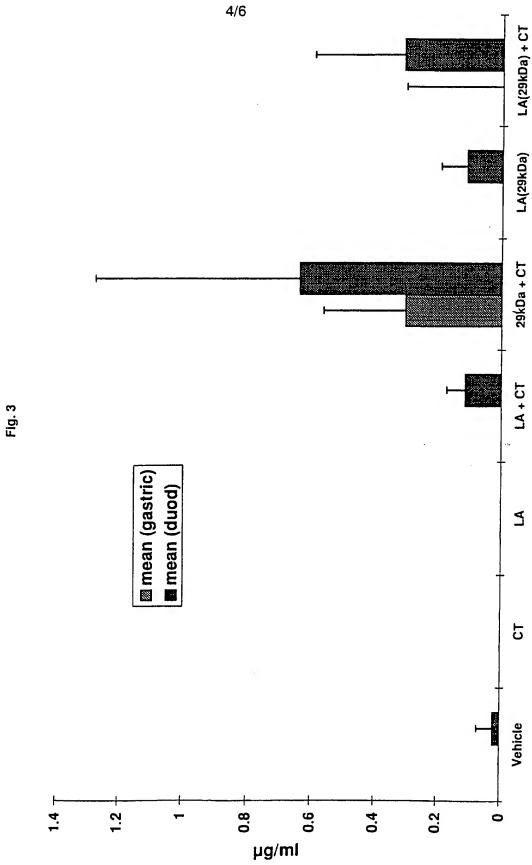
23. A method according to claim 22, wherein the administration is carried out by an oral, parenteral, rectal or nasal route.

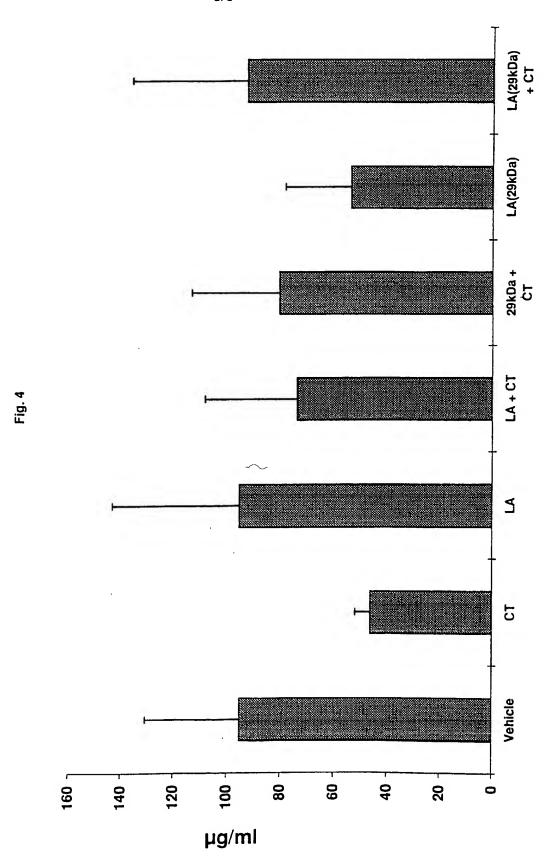


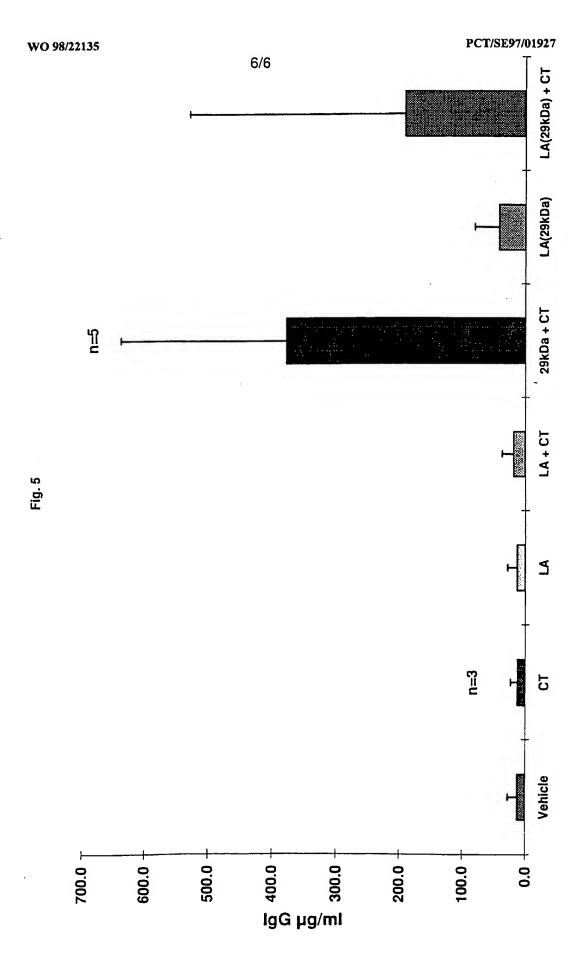
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#### INTERNATIONAL SEARCH REPORT

International application No. PCT/SE 97/01927

### A. CLASSIFICATION OF SUBJECT MATTER IPC6: A61K 39/106, A61K 9/127 // C07K 14/205 According to International Patent Classification (IPC) or to both national classification and IPC B. FIELDS SEARCHED Minimum documentation searched (classification system followed by classification symbols) IPC6: A61K, C07K Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched SE, DK, FI, NO classes as above Electronic data base consulted during the international search (name of data base and, where practicable, search terms used) WPI, MEDLINE, SCISEARCH, CA, EMBASE, BIOSIS C. DOCUMENTS CONSIDERED TO BE RELEVANT Category\* Citation of document, with indication, where appropriate, of the relevant passages Relevant to claim No. Υ Microbiology, Volume 141, 1995, Catherine J. Luke 1-23 et al, "Identification of a 29 kDa flagellar sheath protein in Helicobacter pylori using a murine monoclonal antibody" page 597 - page 604 Υ WO 9509648 A1 (GOULD-FOGERITE, SUSAN), 1-23 13 April 1995 (13.04.95) Y EMBL, Databas GenBank/DDBJ, accession no. 6 X92502, Jones, A.C. et al: "Gene Cloning of flagellar sheat protein of Helicobacter pylori"; & J. Bacteriol. 175 (3), 674-683 (1993) X Further documents are listed in the continuation of Box C. See patent family annex. later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention Special categories of cited documents: document defining the general state of the art which is not considered to be of particular relevance ertier document but published on or after the international filing date "X" document of particular relevance: the claimed invention cannot be considered novel or cannot be considered to involve an inventive document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other step when the document is taken alone special reason (as specified) "Y" document of particular relevance: the claimed invention cannot be "O" document referring to an oral disclosure, use, exhibition or other considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art document published prior to the international filing date but later than the priority date claimed "&" document member of the same patent family Date of the actual completion of the international search Date of mailing of the international search report 24 February 1998 Name and mailing address of the ISA/ Authorized officer Swedish Patent Office Box 5055, S-102 42 STOCKHOLM Patrick Andersson Facsimile No. +46 8 666 02 86 Telephone No. +46 8 782 25 00

# INTERNATIONAL SEARCH REPORT

International application No.
PCT/SE 97/01927

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C (Continue	ation). DOCUMENTS CONSIDERED TO BE RELEVANT			
Category*	Citation of document, with indication, where appropriate, of the rele	vant passages	Relevant to claim No	
A	Gut, Volume 37, No 1, 1995, A Jones et al, "G cloning of a flagellar sheath protein of Helicobacter Pylori", page A63	1-23		
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Information on patent family members

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